

Appl. No. : 09/743,905  
Filed : April 24, 2001

### **AMENDMENTS TO THE SPECIFICATION**

**Please amend line 13 on page 9 of the Specification as filed as follows:**

SEQ ID NO: [[2]] 3

**Please amend line 1 on page 10 of the Specification as filed as follows:**

SEQ ID NO: [[3]] 5

**Please amend paragraph on page 10, lines 23-28 as follows:**

In the following description, the various modified BNYVV TGB-3 sequences will be hereafter called "P15 mutants", identified by the following reference: BNP15-Ala1, corresponding to SEQ ID NO: 1 and SEQ ID NO: 2, BNP15-Ala4 corresponding to SEQ ID NO: [[2]] 3 and SEQ ID NO: 4, BNP15-Asp9, corresponding to SEQ ID NO: [[3]] 5 and SEQ ID NO: 6.

**Please amend paragraph on page 10, lines 29-33 as follows:**

The nucleotidic and corresponding amino-acid sequences of SEQ ID NO: 1, SEQ ID NO: [[2]] 3 and SEQ ID NO: [[3]] 5 can be compared to SEQ ID NOS: [[4]] 7 and 8, which [[is]]are the sequences of the wild type P15 nucleotidic and amino-acid sequence already described (17).

**Please amend paragraph on page 11, line 22 through page 12, line 6 as follows:**

Preferably, said method is used for inducing a resistance to the BNYVV into a sugar beet plant or a sugar beet cell. Said method comprises the following steps:

- preparing a nucleic acid construct comprising a modified nucleic acid sequence obtained by the method according to the invention, preferably preparing a nucleic acid construct comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: [[2]] 3 or SEQ ID NO [[3]] 5, being operably linked to one or more regulatory sequences active into a plant,
- transforming the sugar beet plant cell with the nucleic acid construct, and
- possibly regenerating the transgenic sugar beet plant from the transformed sugar beet plant cell.

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**Please amend paragraph on page 11, lines 16-20 as follows:**

Preferably, said modified nucleic acid sequence is selected from the group consisting of SEQ ID NO: 1, SEQ ID NO [[2]] 3 and SEQ ID NO [[3]] 5, being operably linked to one or more regulatory sequences active into a plant or a plant cell.